

2007-09-27 12090-05CIP2 Sequence Listing.ST25.txt
SEQUENCE LISTING

<110> CHUN, Keun Ho
HWANG, Hyun Jin

<120> TARGET DETECTION SYSTEM HAVING A CONFORMATIONALLY
SENSITIVE PROBE
COMPRISING A NUCLEIC ACID BASED SIGNAL TRANSDUCER

<130> 12090-05CIP2

<140> US 10/684,346
<141> 2003-10-11

<150> US 60/417,864
<151> 2002-10-11

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<313> (88)..(95)

<400> 76

His Ala His Lys Leu Arg Val Asp

1 5

<210> 77

<211> 8

<212> PRT

<213> Homo sapiens

<220>

<221> MISC_FEATURE

<222> (1)..(8)

<223> Falcilysin cleavage site

<400> 77

Trp Thr Gln Arg Phe Phe Glu Ser
1 5

<210> 78

<211> 8

<212> PRT

<213> Homo sapiens

<220>

<221> MISC_FEATURE

<222> (1)..(8)

<223> Falcilysin cleavage site

<400> 78

Ala Phe Ser Asp Gly Leu Ala His
1 5

<210> 79

<211> 8

<212> PRT

<213> Homo sapiens

<220>

<221> MISC_FEATURE

<222> (1)..(8)

<223> Falcilysin cleavage site

<400> 79

Leu Ala His Leu Asp Asn Leu Lys
1 5

<210> 80

<211> 8

<212> PRT

<213> Homo sapiens

<220>
<221> MISC_FEATURE
<222> (1)..(8)
<223> Falcilysin cleavage site

<400> 80

Ala Tyr Gln Lys Val Val Ala Gly
1 5

<210> 81
<211> 8
<212> PRT
<213> Homo sapiens

<220>
<221> MISC_FEATURE
<222> (1)..(8)
<223> Cathepsin cleavage site

<300>
<308> GenBank/NM_000558
<309> 2003-10-04
<313> (27)..(34)

<400> 81

Ala Glu Ala Leu Glu Arg Met Phe
1 5

<210> 82
<211> 8
<212> PRT
<213> Homo sapiens

<220>
<221> MISC_FEATURE
<222> (1)..(8)
<223> Cathepsin cleavage site

<300>

<308> GenBank/NM_000558

<309> 2003-10-04

<313> (34)..(41)

<400> 82

Phe Leu Ser Phe Pro Thr Thr Lys

1 5

<210> 83

<211> 8

<212> PRT

<213> Homo sapiens

<220>

<221> MISC_FEATURE

<222> (1)..(8)

<223> Cathepsin cleavage site

<400> 83

Thr Pro Glu Glu Lys Ala Ser Val

1 5

<210> 84

<211> 8

<212> PRT

<213> Homo sapiens

<220>

<221> MISC_FEATURE

<222> (1)..(8)

<223> Cathepsin cleavage site

<400> 84

Val Thr Ala Leu Trp Glu Lys Val

1 5

<210> 85

<211> 8

<212> PRT

<213> Homo sapiens

<220>

<221> MISC_FEATURE

<222> (1)..(8)

<223> Cathepsin cleavage site

<400> 85

Leu Gly Arg Leu Leu Leu Val Val
1 5

<210> 86

<211> 11

<212> PRT

<213> Artificial

<220>

<223> Synthetic sequence

<220>

<221> MISC_FEATURE

<222> (1)..(11)

<223> cAMP-dependent protein Kinase phosphorylation site

<400> 86

Tyr Leu Arg Arg Ala Ser Leu Ala Gln Leu Thr
1 5 10

<210> 87

<211> 8

<212> PRT

<213> Homo sapiens

<220>

<221> MISC_FEATURE

<222> (1)..(8)

<223> cAMP-dependent protein Kinase phosphorylation site

<400> 87

Phe Arg Arg Leu Ser Ile Ser Thr
1 5

<210> 88

<211> 11

<212> PRT

<213> Bos taurus

<220>

<221> MISC_FEATURE

<222> (1)..(11)

<223> cAMP-dependent protein Kinase phosphorylation site

<400> 88

Ala Gly Ala Arg Arg Lys Ala Ser Gly Pro Pro
1 5 10

<210> 89

<211> 8

<212> PRT

<213> Homo sapiens

<220>

<221> MISC_FEATURE

<222> (1)..(8)

<223> cAMP-dependent protein Kinase phosphorylation site

<400> 89

Gly Arg Gly Leu Ser Leu Ser Arg
1 5

<210> 90

<211> 11

<212> PRT

<213> Oryctolagus cuniculus

<220>
 <221> MISC_FEATURE
 <222> (1)..(11)
 <223> Casein Kinase I phosphorylation site; Ser (location:4)
 phosphorylated

<400> 90

Arg Thr Leu Ser Val Ser Ser Leu Pro Gly Leu
 1 5 10

<210> 91
 <211> 10
 <212> PRT
 <213> Bos taurus

<220>
 <221> MISC_FEATURE
 <222> (1)..(10)
 <223> Casein Kinase I phosphorylation site; Ser (location:4 and 6)
 phosphorylated

<400> 91

Asp Ile Gly Ser Glu Ser Thr Glu Asp Gln
 1 5 10

<210> 92
 <211> 10
 <212> PRT
 <213> Bos taurus

<220>
 <221> MISC_FEATURE
 <222> (1)..(10)
 <223> Casein Kinase II phosphorylation site

<400> 92

Ala Asp Ser Glu Ser Glu Asp Glu Glu Asp
 1 5 10

<210> 93
<211> 11
<212> PRT
<213> Homo sapiens

<220>
<221> MISC_FEATURE
<222> (1)..(11)
<223> Casein Kinase II phosphorylation site

<400> 93

Leu Glu Ser Glu Glu Glu Gly Val Pro Ser Thr
1 5 10

<210> 94
<211> 11
<212> PRT
<213> Homo sapiens

<220>
<221> MISC_FEATURE
<222> (1)..(11)
<223> Casein Kinase II phosphorylation site

<400> 94

Glu Asp Asn Ser Glu Asp Glu Ile Ser Asn Leu
1 5 10

<210> 95
<211> 9
<212> PRT
<213> Homo sapiens

<220>
<221> MISC_FEATURE
<222> (1)..(9)
<223> Glycogen Synthase Kinase 3 phosphorylation site: Ser (location:9)
phosphorylated

<400> 95

Ser Val Pro Pro Ser Pro Ser Leu Ser
1 5

<210> 96

<211> 9

<212> PRT

<213> Homo sapiens

<220>

<221> MISC_FEATURE

<222> (1)..(9)

<223> Glycogen Synthase Kinase 3 phosphorylation site: Ser (location: 5
and 9) phosphorylated

<400> 96

Ser Val Pro Pro Ser Pro Ser Leu Ser
1 5

<210> 97

<211> 7

<212> PRT

<213> Aspergillus fumigatus

<220>

<221> MISC_FEATURE

<222> (1)..(7)

<223> Cdc2 protein Kinase phosphorylation site

<400> 97

Pro Ala Lys Thr Pro Val Lys
1 5

<210> 98

<211> 10

<212> PRT

<213> Simian virus 40

<220>
<221> MISC_FEATURE
<222> (1)..(10)
<223> Cdc2 protein Kinase phosphorylation site

<400> 98

His Ser Thr Pro Pro Lys Lys Lys Arg Lys
1 5 10

<210> 99
<211> 11
<212> PRT
<213> Homo sapiens

<220>
<221> MISC_FEATURE
<222> (1)..(11)
<223> Calmodulin-dependent protein Kinase II phosphorylation site

<400> 99

Asn Tyr Leu Arg Arg Arg Leu Ser Asp Ser Asn
1 5 10

<210> 100
<211> 10
<212> PRT
<213> Homo sapiens

<220>
<221> MISC_FEATURE
<222> (1)..(10)
<223> Calmodulin-dependent protein Kinase II phosphorylation site

<400> 100

Lys Met Ala Arg Val Phe Ser Val Leu Arg
1 5 10

<210> 101
<211> 13
<212> PRT
<213> Artificial

<220>
<223> Synthetic sequence

<220>
<221> MISC_FEATURE
<222> (1)..(13)
<223> Insulin receptor phosphorylation site

<400> 101

Arg Arg Leu Ile Glu Asp Ala Glu Tyr Ala Ala Arg Gly
1 5 10

<210> 102
<211> 4
<212> PRT
<213> Homo sapiens

<220>
<221> MISC_FEATURE
<222> (1)..(4)
<223> Mitogen-activated protein Kinase (Extracellular Signal-regulated
Kinase) phosphorylation site

<400> 102

Pro Leu Ser Pro
1

<210> 103
<211> 4
<212> PRT
<213> Xenopus laevis

<220>

<221> MISC_FEATURE
 <222> (1)..(4)
 <223> Mitogen-activated protein Kinase (Extracellular Signal-regulated Kinase) phosphorylation site

<400> 103

Pro Ser Ser Pro
 1

<210> 104
 <211> 4
 <212> PRT
 <213> Homo sapiens

<220>
 <221> MISC_FEATURE
 <222> (1)..(4)
 <223> Mitogen-activated protein Kinase (Extracellular Signal-regulated Kinase) phosphorylation site

<400> 104

Val Leu Ser Pro
 1

<210> 105
 <211> 21
 <212> PRT
 <213> Homo sapiens

<220>
 <221> MISC_FEATURE
 <222> (1)..(21)
 <223> Mitogen-activated protein Kinase (Extracellular Signal-regulated Kinase) phosphorylation site

<400> 105

Lys Arg Glu Leu Val Glu Pro Leu Thr Pro Ser Gly Glu Ala Pro Asn
 1 5 10 15

Gln Ala Leu Leu Arg
20

<210> 106
<211> 11
<212> PRT
<213> Homo sapiens

<220>
<221> MISC_FEATURE
<222> (1)..(11)
<223> cGMP-dependent protein Kinase phosphorylation site

<400> 106

Gly Lys Lys Arg Lys Arg Ser Arg Lys Glu Ser
1 5 10

<210> 107
<211> 8
<212> PRT
<213> Bos taurus

<220>
<221> MISC_FEATURE
<222> (1)..(8)
<223> cGMP-dependent protein Kinase phosphorylation site

<400> 107

Phe Arg Arg Leu Ser Ile Ser Thr
1 5

<210> 108
<211> 7
<212> PRT
<213> Artificial

<220>
<223> Synthetic sequence

<220>
<221> MISC_FEATURE
<222> (1)..(7)
<223> cGMP-dependent protein Kinase phosphorylation site

<400> 108

Arg Lys Arg Ser Arg Ala Glu
1 5

<210> 109
<211> 12
<212> PRT
<213> Homo sapiens

<220>
<221> MISC_FEATURE
<222> (1)..(12)
<223> Phosphorylase Kinase phosphorylation site

<400> 109

Asp Gln Glu Lys Arg Lys Gln Ile Ser Val Arg Gly
1 5 10

<210> 110
<211> 10
<212> PRT
<213> Artificial

<220>
<223> Synthetic sequence

<220>
<221> MISC_FEATURE
<222> (1)..(10)
<223> Phosphorylase Kinase phosphorylation site

<400> 110

Pro Leu Ser Arg Thr Leu Ser Val Ser Ser

1 5 10

<210> 111

<211> 9

<212> PRT

<213> Homo sapiens

<220>

<221> MISC_FEATURE

<222> (1)..(9)

<223> Protein Kinase C phosphorylation site

<400> 111

His Glu Gly Thr His Ser Thr Lys Arg

1 5

<210> 112

<211> 10

<212> PRT

<213> Artificial

<220>

<223> Synthetic sequence

<220>

<221> MISC_FEATURE

<222> (1)..(10)

<223> Protein Kinase C phosphorylation site

<400> 112

Pro Leu Ser Arg Thr Leu Ser Val Ser Ser

1 5 10

<210> 113

<211> 11

<212> PRT

<213> Artificial

<220>

<223> Synthetic sequence

<220>

<221> MISC_FEATURE

<222> (1)..(11)

<223> Protein Kinase C phosphorylation site

<400> 113

Gln Lys Arg Pro Ser Gln Arg Ser Lys Tyr Leu
1 5 10

<210> 114

<211> 12

<212> PRT

<213> Artificial

<220>

<223> Synthetic sequence

<220>

<221> MISC_FEATURE

<222> (1)..(12)

<223> Protein Kinase C phosphorylation site

<400> 114

Pro Leu Ser Arg Thr Leu Ser Val Ala Ala Lys Lys
1 5 10

<210> 115

<211> 7

<212> PRT

<213> Artificial

<220>

<223> Synthetic sequence

<220>

<221> MISC_FEATURE

<222> (1)..(7)

<223> Protein Kinase C phosphorylation site

<400> 115

Leu Lys Phe Ser Lys Lys Phe
1 5

<210> 116

<211> 8

<212> PRT

<213> Homo sapiens

<220>

<221> MISC_FEATURE

<222> (1)..(8)

<223> Protein Kinase C phosphorylation site

<400> 116

Arg Lys Arg Thr Leu Arg Arg Leu
1 5

<210> 117

<211> 21

<212> PRT

<213> Artificial

<220>

<223> Synthetic sequence

<220>

<221> MISC_FEATURE

<222> (1)..(21)

<223> p34 cdc2 protein Kinase phosphorylation site

<400> 117

Ala Lys Ala Gln His Ala Thr Pro Pro Lys Lys Lys Arg Lys Val Glu
1 5 10 15

Asp Pro Lys Asp Phe

20

<210> 118

<211> 9

<212> PRT

<213> Artificial

<220>

<223> Synthetic sequence

<220>

<221> MISC_FEATURE

<222> (1)..(9)

<223> Meiosis-activated myelin basic protein Kinase phosphorylation
site

<400> 118

Ala Pro Arg Thr Pro Gly Gly Arg Arg

1

5

<210> 119

<211> 11

<212> PRT

<213> Artificial

<220>

<223> Synthetic sequence

<220>

<221> MISC_FEATURE

<222> (1)..(11)

<223> Smooth Muscle Myosin Light Chain Kinase phosphorylation site

<400> 119

Lys Lys Arg Ala Arg Thr Ser Asn Val Phe Ala

1

5

10

<210> 120

<211> 11
<212> PRT
<213> Artificial

<220>
<223> Synthetic sequence

<220>
<221> MISC_FEATURE
<222> (1)..(11)
<223> Epidermal Growth Factor Receptor Kinase phosphorylation site

<400> 120

Arg Glu Asn Ala Glu Tyr Leu Arg Val Ala Pro
1 5 10

<210> 121
<211> 10
<212> PRT
<213> Homo sapiens

<220>
<221> MISC_FEATURE
<222> (1)..(10)
<223> Epidermal Growth Factor Receptor Kinase phosphorylation site

<400> 121

Ala Glu Pro Asp Tyr Gly Ala Leu Tyr Glu
1 5 10

<210> 122
<211> 5
<212> PRT
<213> Artificial

<220>
<223> Synthetic sequence

<220>

<221> MISC_FEATURE

<222> (1)..(5)

<223> Protein Tyrosine Kinase pp60c-src phosphorylation site

<400> 122

Ile Tyr Gly Glu Phe

1 5

<210> 123

<211> 52

<212> DNA

<213> Artificial

<220>

<223> Synthetic Sequence

<220>

<221> modified_base

<222> (39)..(39)

<223> The 39th nucleotide t is linked to biotin by a linker.

<400> 123

atggaagtat atggaagtat atggaagtat tcgtgggggtt ttgcagtcgt ag 52

<210> 124

<211> 14

<212> DNA

<213> Artificial

<220>

<223> Synthetic Sequence

<220>

<221> modified_base

<222> (1)..(1)

<223> The first nucleotide g is linked to fluorescein by a linker.

<400> 124

gactgcaaaa cccc 14

<210> 125
 <211> 52
 <212> DNA
 <213> Artificial

<220>
 <223> Synthetic Sequence

<220>
 <221> misc_feature
 <222> (39)..(39)
 <223> The 39th nucleotide n is an abasic nucleotide,
 6-amino-2-hydroxymethyl hexanol linked to biotin.

<400> 125
 atggaagtat atggaagtat atggaagtat tcgtggggnt ttgcagtcgt ag 52

<210> 126
 <211> 16
 <212> DNA
 <213> Artificial

<220>
 <223> Synthetic Sequence

<220>
 <221> modified_base
 <222> (1)..(1)
 <223> The first nucleotide g is linked to fluorescein by a linker.

<400> 126
 gactgcaaaa ccccac 16

<210> 127
 <211> 36
 <212> DNA
 <213> Artificial

<220>
 <223> Synthetic Sequence

<220>
 <221> modified_base
 <222> (1)..(1)
 <223> The first nucleotide g is linked to fluorescein by a linker.

<220>
 <221> stem_loop
 <222> (1)..(36)

<220>
 <221> modified_base
 <222> (34)..(34)
 <223> The 34th nucleotide t is linked to biotin by a linker.

<220>
 <221> modified_base
 <222> (36)..(36)
 <223> The last (36th) nucleotide c is linked to DABCYL
 (4-(4'-dimethylaminophenylazo)benzoic acid) by a linker.

<400> 127
 gcagcctagg aaacaccaaa gatgatattt ggctgc 36

<210> 128
 <211> 38
 <212> DNA
 <213> Artificial

<220>
 <223> Synthetic Sequence

<220>
 <221> modified_base
 <222> (1)..(1)
 <223> The first nucleotide g is linked to fluorescein by a linker.

<220>
 <221> stem_loop
 <222> (1)..(38)

<220>
 <221> modified_base
 <222> (6)..(6)
 <223> The 6th nucleotide t is linked to biotin by a linker.

<220>
 <221> modified_base
 <222> (36)..(36)
 <223> The 36th nucleotide t is linked to biotin by a linker.

<220>
 <221> modified_base
 <222> (38)..(38)
 <223> The last (38th) nucleotide c is linked to DABCYL
 (4-(4'-dimethylaminophenylazo)benzoic acid) by a linker.

<400> 128
 gcagctctag gaaacaccaa agatgatatt tgagctgc 38

<210> 129
 <211> 30
 <212> DNA
 <213> Artificial

<220>
 <223> Synthetic Sequence

<400> 129
 aaatatcatc ttggtgttt cctaggctgc 30

<210> 130
 <211> 14
 <212> DNA
 <213> Artificial

<220>
 <223> Synthetic Sequence

<400> 130
 gactgcaaaa cccc 14

<210> 131
 <211> 12
 <212> DNA
 <213> Artificial

<220>

<223> Synthetic Sequence

<220>

<221> modified_base

<222> (1)..(1)

<223> The first nucleotide c is linked to fluorescein by a linker.

<400> 131

ctacgactgc aa

12

<210> 132

<211> 52

<212> DNA

<213> Artificial

<220>

<223> Synthetic Sequence

<400> 132

atggaagtat atggaagtat atggaagtat tcgtgggggtt ttgcagtcgt ag

52

<210> 133

<211> 14

<212> DNA

<213> Artificial

<220>

<223> Synthetic Sequence

<220>

<221> modified_base

<222> (4)..(4)

<223> The 4th nucleotide t is linked to biotin by a linker.

<400> 133

gactgcaaaa cccc

14

<210> 134

<211> 16

<212> DNA

<213> Artificial

<220>
<223> Synthetic Sequence

<400> 134
gactgcaaaa ccccac 16

<210> 135
<211> 52
<212> DNA
<213> Artificial

<220>
<223> Synthetic Sequence

<220>
<221> modified_base
<222> (1)..(1)
<223> The first nucleotide a is linked to biotin by a linker.

<400> 135
atggaagtat atggaagtat atggaagtat tcgtgggggtt ttgcagtcgt ag 52

<210> 136
<211> 16
<212> DNA
<213> Artificial

<220>
<223> Synthetic Sequence

<220>
<221> modified_base
<222> (4)..(4)
<223> The 4th nucleotide t is linked to biotin by a linker.

<400> 136
gactgcaaaa ccccac 16

<210> 137
<211> 52

<212> DNA

<213> Artificial

<220>

<223> Synthetic Sequence

<220>

<221> modified_base

<222> (1)..(1)

<223> The first nucleotide a is linked to biotin by a linker.

<220>

<221> misc_feature

<222> (39)..(39)

<223> The 39th nucleotide n is an abasic nucleotide,
6-amino-2-hydroxymethyl hexanol linked to digoxigenin.

<400> 137

atggaagtat atggaagtat atggaagtat tcgtggggnt ttgcagtcgt ag

52

<210> 138

<211> 14

<212> DNA

<213> Artificial

<220>

<223> Synthetic sequence

<220>

<221> stem_loop

<222> (1)..(14)

<220>

<221> modified_base

<222> (12)..(12)

<223> The 12nd nucleotide t is linked to biotin by a linker.

<400> 138

gcaggactac ctgc

14

<210> 139

<211> 16

<212> DNA

<213> Artificial

<220>

<223> Synthetic sequence

<220>

<221> stem_loop

<222> (1)..(16)

<220>

<221> modified_base

<222> (14)..(14)

<223> The 14th nucleotide t is linked to biotin by a linker.

<400> 139

gcaggacttt acctgc

16

<210> 140

<211> 18

<212> DNA

<213> Artificial

<220>

<223> Synthetic sequence

<220>

<221> stem_loop

<222> (1)..(18)

<220>

<221> modified_base

<222> (16)..(16)

<223> The 16th nucleotide t is linked to biotin by a linker.

<400> 140

gcaggactca ttacctgc

18

<210> 141

<211> 25

<212> DNA

<213> Artificial

<220>
<223> Synthetic sequence

<220>
<221> stem_loop
<222> (1)..(25)

<220>
<221> modified_base
<222> (23)..(23)
<223> The 23rd nucleotide t is linked to biotin by a linker.

<400> 141
gcaggatact cattaccata cctgc 25

<210> 142
<211> 35
<212> DNA
<213> Artificial

<220>
<223> Synthetic sequence

<220>
<221> stem_loop
<222> (1)..(35)

<220>
<221> modified_base
<222> (33)..(33)
<223> The 33rd nucleotide t is linked to biotin by a linker.

<400> 142
gcaggatact cattagcgac gaacaccata cctgc 35

<210> 143
<211> 45
<212> DNA
<213> Artificial

<220>

<223> Synthetic sequence

<220>

<221> stem_loop

<222> (1)..(45)

<220>

<221> modified_base

<222> (43)..(43)

<223> The 43rd nucleotide t is linked to biotin by a linker.

<400> 143

gcaggatact tagaccaaca cattagcgac gaacaccata cctgc 45

<210> 144

<211> 25

<212> DNA

<213> Artificial

<220>

<223> Synthetic sequence

<220>

<221> stem_loop

<222> (1)..(25)

<220>

<221> modified_base

<222> (23)..(23)

<223> The 23rd nucleotide t is linked to biotin by a linker.

<400> 144

cgaccatcct cattaccata ggtagc 25

<210> 145

<211> 25

<212> DNA

<213> Artificial

<220>

<223> Synthetic sequence

<220>
 <221> stem_loop
 <222> (1)..(25)

 <220>
 <221> modified_base
 <222> (23)..(23)
 <223> The 23rd nucleotide t is linked to biotin by a linker.

<400> 145
 gcagcatcct cattacccta gctgc 25

<210> 146
 <211> 25
 <212> DNA
 <213> Artificial

<220>
 <223> Synthetic sequence

<220>
 <221> stem_loop
 <222> (1)..(25)

 <220>
 <221> modified_base
 <222> (23)..(23)
 <223> The 23rd nucleotide t is linked to biotin by a linker.

<400> 146
 cgacgatcct cattaccata cgtcg 25

<210> 147
 <211> 25
 <212> DNA
 <213> Artificial

<220>
 <223> Synthetic sequence

<220>

<221> stem_loop

<222> (1)..(25)

<220>

<221> modified_base

<222> (23)..(23)

<223> The 23rd nucleotide t is linked to biotin by a linker.

<400> 147

ggaggataat cattaccata cctcc

25

<210> 148

<211> 25

<212> DNA

<213> Artificial

<220>

<223> Synthetic sequence

<220>

<221> stem_loop

<222> (1)..(25)

<220>

<221> modified_base

<222> (23)..(23)

<223> The 23rd nucleotide t is linked to biotin by a linker.

<400> 148

ccaccatact cattacccta ggtgg

25

<210> 149

<211> 23

<212> DNA

<213> Artificial

<220>

<223> Synthetic sequence

<220>

<221> stem_loop

<222> (1)..(23)

<220>
 <221> modified_base
 <222> (21)..(21)
 <223> The 21st nucleotide t is linked to biotin by a linker.

<400> 149
 gcagatactc attaccatac tgc 23

<210> 150
 <211> 25
 <212> DNA
 <213> Artificial

<220>
 <223> Synthetic sequence

<220>
 <221> stem_loop
 <222> (1)..(25)

<220>
 <221> modified_base
 <222> (23)..(23)
 <223> The 23rd nucleotide t is linked to biotin by a linker.

<400> 150
 gcaggatact gcttaccata cctgc 25

<210> 151
 <211> 25
 <212> DNA
 <213> Artificial

<220>
 <223> Synthetic sequence

<220>
 <221> stem_loop
 <222> (1)..(25)

<220>

<221> modified_base
 <222> (23)..(23)
 <223> The 23rd nucleotide t is linked to biotin by a linker.

<400> 151
 gcaggactct cattacactg cctgc 25

<210> 152
 <211> 25
 <212> DNA
 <213> Artificial

<220>
 <223> Synthetic sequence

<220>
 <221> modified_base
 <222> (25)..(25)
 <223> The 25th nucleotide t is linked to biotin by a linker.

<400> 152
 agcgcatcct cattacccta gcgct 25

<210> 153
 <211> 25
 <212> DNA
 <213> Artificial

<220>
 <223> Synthetic sequence

<220>
 <221> stem_loop
 <222> (1)..(25)

<220>
 <221> modified_base
 <222> (21)..(21)
 <223> The 21st nucleotide t is linked to biotin by a linker.

<400> 153
 gcgcaatcct cattacccta tgcgc 25

<210> 154
 <211> 25
 <212> DNA
 <213> Artificial

<220>
 <223> Synthetic sequence

<220>
 <221> stem_loop
 <222> (1)..(25)

<220>
 <221> modified_base
 <222> (19)..(19)
 <223> The 19th nucleotide t is linked to biotin by a linker.

<400> 154
 gcagcatcct cattacccta gctgc 25

<210> 155
 <211> 25
 <212> DNA
 <213> Artificial

<220>
 <223> Synthetic sequence

<220>
 <221> stem_loop
 <222> (1)..(25)

<220>
 <221> modified_base
 <222> (13)..(13)
 <223> The 13rd nucleotide t is linked to biotin by a linker.

<400> 155
 gcagcatcct cattacccta gctgc 25

<210> 156

<211> 25

<212> DNA

<213> Artificial

<220>

<223> Synthetic sequence

<220>

<221> stem_loop

<222> (1)..(25)

<220>

<221> modified_base

<222> (10)..(10)

<223> The 10th nucleotide t is linked to biotin by a linker.

<400> 156

gcagcatcct cattacccta gctgc

25

<210> 157

<211> 25

<212> DNA

<213> Artificial

<220>

<223> Synthetic sequence

<220>

<221> modified_base

<222> (1)..(1)

<223> The first nucleotide g is linked to fluorescein by a linker.

<220>

<221> stem_loop

<222> (1)..(25)

<220>

<221> modified_base

<222> (25)..(25)

<223> The last (25th) nucleotide c is linked to DABCYL
(4-(4'-dimethylaminophenylazo)benzoic acid) by a linker.

<400> 157
gcagctagga gtaatgggat gctgc 25

<210> 158
<211> 15
<212> DNA
<213> Artificial

<220>
<223> Synthetic sequence

<220>
<221> modified_base
<222> (11)..(11)
<223> The 11st nucleotide t is linked to biotin by a linker.

<400> 158
atccattac tccta 15

<210> 159
<211> 13
<212> DNA
<213> Artificial

<220>
<223> Synthetic sequence

<220>
<221> modified_base
<222> (11)..(11)
<223> The 11st nucleotide t is linked to biotin by a linker.

<400> 159
atccattac tcc 13

<210> 160
<211> 15
<212> DNA
<213> Artificial

<220>

<223> Synthetic sequence

<400> 160

tagggtaatg aggat

15

<210> 161

<211> 25

<212> DNA

<213> Artificial

<220>

<223> Synthetic sequence

<220>

<221> stem_loop

<222> (1)..(25)

<220>

<221> modified_base

<222> (23)..(23)

<223> The 23rd nucleotide t is linked to carboxyl group by a linker.

<400> 161

gcagcatcct cattacccta gctgc

25

<210> 162

<211> 25

<212> DNA

<213> Artificial

<220>

<223> Synthetic sequence

<220>

<221> stem_loop

<222> (1)..(25)

<220>

<221> modified_base

<222> (23)..(23)

<223> The 23rd nucleotide t is linked to amine group by a linker.

<400> 162

gcagcatcct cattacccta gctgc

25

<210> 163

<211> 7

<212> PRT

<213> Artificial

<220>

<223> Synthetic sequence

<220>

<221> MISC_FEATURE

<222> (1)..(7)

<223> Protein Kinase C phosphorylation site

<400> 163

Lys Arg Thr Leu Arg Arg Cys

1 5

<210> 164

<211> 6

<212> PRT

<213> Homo sapiens

<220>

<221> MISC_FEATURE

<222> (1)..(6)

<223> Protein Kinase C phosphorylation site

<400> 164

Lys Arg Thr Leu Arg Arg

1 5

<210> 165

<211> 25

<212> DNA

<213> Artificial

<220>

<223> Synthetic sequence

<220>

<221> stem_loop

<222> (1)..(25)

<220>

<221> modified_base

<222> (23)..(23)

<223> The 23rd nucleotide t is linked to phosphorylated heptapeptide, KRpTLRRC, by a linker.

<400> 165

gcagcatcct cattacccta gctgc

25

<210> 166

<211> 7

<212> PRT

<213> Artificial

<220>

<223> Synthetic sequence

<220>

<221> MISC_FEATURE

<222> (3)..(3)

<223> The 3rd amino acid T is phosphorylated.

<400> 166

Lys Arg Thr Leu Arg Arg Cys

1

5

<210> 167

<211> 25

<212> DNA

<213> Artificial

<220>

<223> Synthetic sequence

<220>

<221> stem_loop

<222> (1)..(25)

<220>

<221> modified_base

<222> (23)..(23)

<223> The 23rd nucleotide t is linked to the heptapeptide, KRpTLRRC, by
a linker.

<400> 167

gcagcatcct cattacccta gctgc

25